amino acid sequence alignment between SEQ ID NO: 4 (also identified as "C-type Lectin Receptor-like) and mouse macrophage C-type lectin, a type II transmembrane protein ("macrophage C-type lectin") (SEQ ID NO: 8), indicating that the two sequences share 55% overall similarity and 39% identity. Figure 2 shows the BLASTP amino acid sequence alignment between SEQ ID NO: 4 (also identified as "C-type Lectin Receptor-like) and human dendritic cell immunoreceptor ("dendritic cell immunoreceptor") (SEQ ID NO: 9), indicating that the two sequences share 49% identity and 69% overall similarity. Figure 3 shows the BLASTP amino acid sequence alignment between SEQ ID NO: 4 (also identified as "C-type Lectin Receptor-like) and human C-type lectin DDB27 ("DDB27") (SEQ ID NO: 10), indicating that the two sequences share 49% identity and 69% overall similarity. Figure 4 shows the BLASTP amino acid sequence alignment between SEQ ID NO: 4 (also identified as "C-type Lectin Receptor-like) and mouse C-type (calcium dependent, carbohydrate recognition domain) lectin, superfamily member 6 ("mouse C-type") (SEQ ID NO: 11), indicating that the two sequences share 44% identity and 62% overall similarity. The sequences of the present invention are expected to have C-type lectin receptor activity.

Cont A1

2) At page 10, lines 1 through 19, please delete the four paragraphs and replace with the following:

A2

Figure 1 shows the BLASTP amino acid sequence alignment between SEQ ID NO: 4 (also identified as "C-type Lectin Receptor-like) and mouse macrophage C-type lectin, a type II transmembrane protein ("macrophage C-type lectin") (SEQ ID NO: 8), indicating that the two sequences share 55% overall similarity and 39% identity.

Figure 2 shows the BLASTP amino acid sequence alignment between SEQ ID NO: 4 (also identified as "C-type Lectin Receptor-like) and human dendritic cell immunoreceptor ("dendritic cell immunoreceptor") (SEQ ID NO: 9), indicating that the two sequences share 49% identity and 69% overall similarity.

Figure 3 shows the BLASTP amino acid sequence alignment between SEQ ID NO: 4 (also identified as "C-type Lectin Receptor-like) and human C-type lectin DDB27 ("DDB27") (SEQ ID NO: 10), indicating that the two sequences share 49% identity and 69% overall similarity.

Figure 4 shows the BLASTP amino acid sequence alignment between SEQ ID NO: 4 (also identified as "C-type Lectin Receptor-like) and mouse C-type (calcium dependent, carbohydrate recognition domain) lectin, superfamily member 6 ("mouse C-type") (SEQ ID NO: 11), indicating that the two sequences share 44% identity and 62% overall similarity.

In the Sequence Listing.

Please replace the current Sequence Listing with the enclosed substitute Sequence Listing.